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1501 GCAG 1560
1653 GCAG 1712
1561 AGATGAGAGGAG 1620
1713 AGATGAG 1772
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1833 GAG 1892
1741 CCGCATCACCGCTCAATGAGTGAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1800
1893 CCGCATCACCGCTCAATGAGTGAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1952
1801 GAGCGCGAGCTGGCTCTCAATGAGTGAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 1860
1953 GAGCGCGAGCTGGCTCTCAATGAGTGAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2012
1861 GGAAG 1920
2013 GGAAG 2072
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2221 CACTGTCAACAAACAGCTCAGACAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2280
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2382 GGAACAG 2441
2341 AGGCGCACCCACCCAG 2400
2442 AGGCGCACCCACCCAG 2501

336 GCGCCAGCGGCGGAGCCCTCCCTGCTGTCTGAGTTCCAGCCCGGGAATGAACGGTCCCA 395
396 GCGGAGGAGTTTCATGAGTCCCAAGAACTTTGGCTGTATCGCATCATTTCTTGGAGAG 1380
1473 CCGGAGGAGTTTCATGAGTCCCAAGAACTTTGGCTGTATCGCATCATTTCTTGGAGAG 1532
1381 GAAGACAGTGGCTGAGTGGCTCTCTATTATTAATCTGACTAAGAAAGATGAGAACTATAA 1440
1533 GAAGACAGTGGCTGAGTGGCTCTCTATTATTAATCTGACTAAGAAAGATGAGAACTATAA 1592
1441 GAGCTCTGTGAGAGCGGAGCTATCGGCGCGGCGGCAAGAGCCAGAGCAACAAACAGCAGCA 1500
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1501 GCAG 1560
1653 GCAG 1712
1561 AGATGAGAGGAG 1620
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1621 CGACAG 1680
1773 CGACAG 1832
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1981 GAACCTCGATGAGATCTTGAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAGCAG 2040
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2041 GAGGAG 2100
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2101 GAGGAG 2160
2253 GAGGAG 2312
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2221 CACTGTCAACAAACAGCTCAGACAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAG 2280
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2281 GGAACAG 2340
2382 GGAACAG 2441
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2442 AGGCGCACCCACCCAG 2501

4662	AGGGACCGCCAGAGCTCGGGGGGCTCCATTGTGCGCGGGGCCCGCGTCAATGTGCGCTGA	4721
4597	GCTGGGTAAAGCCGCGGCAGAGCCCCCTGACCTATGAGAGACCA CGGGGCA C C C T T T G C G G	4656
4722	GCTGGGTAAAGCCGCGGCAGAGCCCCCTGACCTATGAGAGCA C C C T T T G C G G	4781
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4777	GGCCAAGTCCCGCACAGCACCGTCGCGAGCACACCCACACCCCATCTCGCCCTATGA	4836
4902	GGCCAAGTCCCGCACAGCACCGTCGCGAGCACACCCACACCCCATCTCGCCCTATGA	4961
4837	GCACCTGCTTCGGGGCGTGAGTGGGTGGACCTGTATTCGAGCCACAATCCCTCGGCTT	4896
4962	GCACCTGCTTCGGGGCGTGAGTGGGTGGACCTGTATTCGAGCCACAATCCCTCGGCTT	5021
4897	CGACCCCACTTCATATACCCCGCGCATCCCTCTCGAGCGACCGCTGCCTACTACCTGCC	4956
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5137	CCTCTCGCCCCGAGTCTCTCGCTGGCACTCAACTACGCTCGGGTCCCCGAGGCACTCAT	5196
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5497	GCAGAGCAGCGGACGAGCGGACGCGCGGGGGTGGGGGCGACGACGACGCGCCCGC	5556
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5557	CTCCACTCCCATGCCACACGACCTCGCCCATCTCCCTTCGGAACCCAGGATGCCCTCCA	5616
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Qy	5737	ATTCCCACTGCCACCACTCTGCCCTCTCTCGCAAGCCCACTCTCGATGGGGTCTTACCTTACCCT	5796
Db	5862	ATTCCCACTGCCACCACTCTGCCCTCTCTCGCAAGCCCACTCTCGATGGGGTCTTACCTTACCCT	5921
Qy	5797	CATGGAGCCCGTCTTGTCTGCCCAAGAGAGGCCCGCCCGGGTCCCGCGGCAGAGCGGCCCGG	5856
Db	5922	CATGGAGCCCGTCTTGTCTGCCCAAGAGAGGCCCGCCCGGGTCCCGCGGCAGAGCGGCCCGG	5981
Qy	5857	AGCAGACAACCGGCAATGCCCTTCTCTCGCAAGCCCAAGCCCGCTCGGGGCTTGAGAGCCCGC	5916
Db	5982	AGCAGACAACCGGCAATGCCCTTCTCTCGCAAGCCCAAGCCCGCTCGGGGCTTGAGAGCCCGC	6041
Qy	5917	CTCTCTCCCAAGCAAGGGCTCGAGAGCCCGCCCTCTAGTGCCTCTCTGTCTCTTGCCCAAGC	5976
Db	6042	CTCTCTCCCAAGCAAGGGCTCGAGAGCCCGCCCTCTAGTGCCTCTCTGTCTCTTGCCCAAGC	6101
Qy	5977	CACATCGCCCGCACCCCTCTCGAAGAACCTTCGCACCTCACACGCGCAGCCCGGAGCCCGCC	6036
Db	6102	CACATCGCCCGCACCCCTCTCGAAGAACCTTCGCACCTCACACGCGCAGCCCGGAGCCCGCC	6161
Qy	6037	GGCGCCAATCTGCTCGGCCCTCGGACCCCGCACCGGGAAGAAGCTCAAAAGTAAACCTTTTTC	6096
Db	6162	GGCGCCAATCTGCTCGGCCCTCGGACCCCGCACCGGGAAGAAGCTCAAAAGTAAACCTTTTTC	6221
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Db	6222	CATCTCAGGAACCTGGAACTCGGTTCTCTGGGTTACCAAGGAGAGCTACAGCCCGCGAAGG	6281
Qy	6157	GGTGGAGCCCGTCAAGCCTCTGTGAGCTCACCCAGCTCTGACCCACGACAAGGGGCTTCCCAA	6216
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Qy	6217	GCACTCTGGAAGACTCTGAACAAGAGCCACTCTGAGGGGAGCTGCGGCCCAAGCAGCAGG	6276
Db	6342	GCACTCTGGAAGACTCTGAACAAGAGCCACTCTGAGGGGAGCTGCGGCCCAAGCAGCAGG	6401
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Db	6462	CCAGCCCTCGTCCAGCCCGCTGTCTCCAGACCGGCCCAAGGGGTCAAAAGTCAACAGGGGT	6521
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Db	6522	GGTCAACCTTGGGCCAGCAATCATGTGAGGTATCAACAAGACTACACCCGGGACCAACCC	6581
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7892 CCTTACTCAGGGATGTTTACTGTGTGCTCGGGAAGGAGGAGGAGGAGGCGGGAGGGG 7951

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7897	QY	GATGACCA CGCA CTTCCA CGC CACTGCTCCCGCGAATGCATTTGGAAACCAAGTCTAAA	7956
8012	DB	GATGACCA CGCA CTTCCA CGC CACTGCTCCCGCGAATGCATTTGGAAACCAAGTCTAAA	8071
7957	QY	CTGAGCTCGCAGCCCCCGCGCCCTCCCTCGCCTCCCATCCGCTTAGCGCTCTGGACAG	8016
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8017	QY	ATGGACG CAG GCCCTGTG C CAG CCCCCCAGTCGGTCTCGTTC CGGTCCCA CACAGCTGCCCCA	8076
8132	DB	ATGGACG CAG GCCCTGTG C CAG CCCCCCAGTCGGTCTCGTTC CGGTCCCA CACAGCTGCCCCA	8191
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8257	QY	CTTCAATGAATTAATTCAGATGTTTTACGCAAGGAAGCACTTACCAGTATCTAGCTGC	8316
8372	DB	CTTCAATGAATTAATTCAGATGTTTTACGCAAGGAAGCACTTACCAGTATCTAGCTGC	8431
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8432	DB	TGTGCTTTTGATCTCTGCTTACGTTCAAGAGCGGTGTCAGGCCGACAGTCGGTGAACC	8491
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8492	DB	CATCATCTGCAGGACCAAGGGCGGGGACTGCTCGTCA CGCCCGCGTGTCTCCCTC	8551
8437	QY	CCTCCCTTCCTTGGGCAAGATGAATTCGATGCGTATTCGTGGCGGCCATTTGCGCAGGG	8496
8552	DB	CCTCCCTTCCTTGGGCAAGATGAATTCGATGCGTATTCGTGGCGGCCATTTGCGCAGGG	8611
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8672	DB	AAAAAAAAAAAAAA 8686	

[illegible]

KEYWORDS	Unknown.
SOURCE	Unknown.
ORGANISM	Unknown.

ORGANISM: *Salmonella*
 UNCLASSIFIED.
 (pages 1 to 9053)

REFERENCE	1 (bases 1 to 9053)
AUTHORS	Furness, L.M. and Buchbinder, J.L.
TITLE	Genes expressed in C3A liver cell cultures treated with steroids
JOURNAL	Patent: US 6673549-A 306 06-JAN-2004;
FEATURES	Incyte Corporation; Palo Alto, CA
SOURCE	Location/Qualifiers
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ORIGIN

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